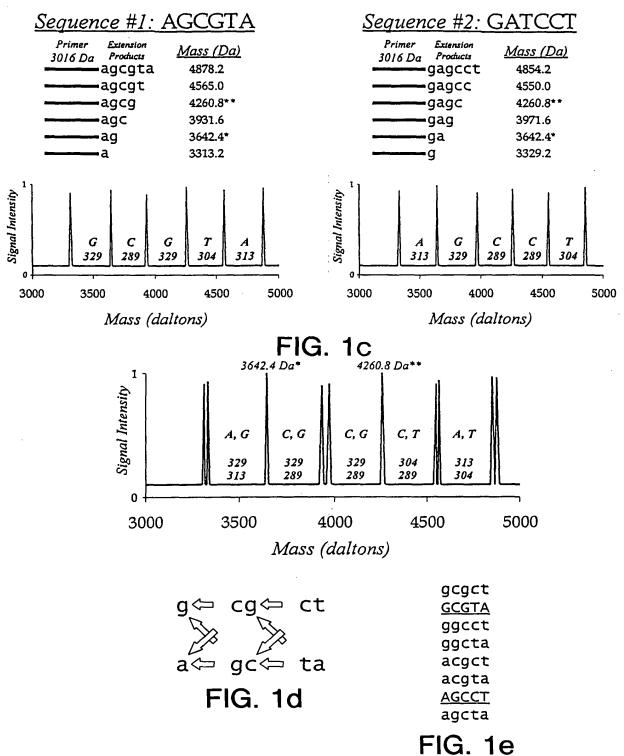
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Sheet I of 8

Title: USE OF NUCLEOTIDE ANALOGS IN THE ANALYSIS OF
OLIGONUCLEOTIDE MIXTURES AND IN HIGHLY
MULTIPLEXED NUCLEIC ACID SEQUENCING
Docket No.: 25491-2408B
ANDICAN: Control of

Applicant: Cantor et al. Filed: August 20, 2003

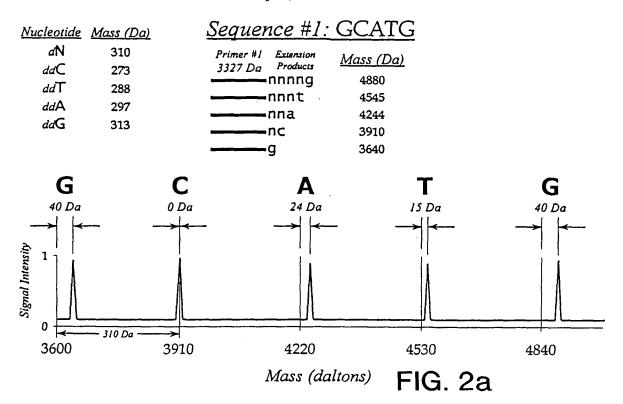


FIG. 1b



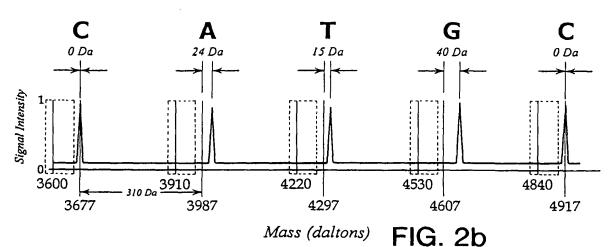
HELLER EHRMAN WHITE & MCAULIFFE LLP
Sheet 2 of 8
Title: USE OP NUCLEOTIDE ANALOGS IN THE ANALYSIS OF
OLIGONUCLEOTIDE MIXTURES AND IN HIGHLY
MULTIPLEXED NUCLEIC ACID SEQUENCING
Docket No.: 25491-2408B

Applicant: Cantor et al. Filed: August 20, 2003



Sequence #2: CATGC

Primer #2 Extension 3404 Da Products	Mass (Da)
nnnnc	4917
nnng	4647
nnt	4312
——na	4011
C	3677



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Sheet 3 of 8

Title: USE OF NUCLECTIBE ANALOGS IN THE ANALYSIS OF
OLIGONUCLECTIBE MIXTURES AND IN HIGHLY
MULTIPLEXED NUCLEIC ACID SEQUENCING
Docket No.: 25491-2408B
Applicant: Cantor et al.
Filed: August 20, 2003

3640

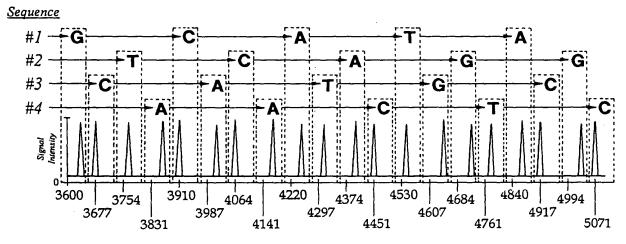
Sequence #1: (GCATA	Sequence #3: C	CATGC
Primer #1 Extension 3327 Da Products	Mass (Da)	Primer #3 Extension 3404 Da Products	Mass (Da)
nnnna	4864	nnnc	4917
nnnt	4545	nnng	4647
nna	4244	nnt	4312
——nc	3910	na	4011

Sequence #2:	ノンハンエ	3
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Sequence #4: AACTC

3677

Primer #2 Extension 3481 Da Products ————————————————————————————————————	<u>Mass (Da)</u> 5034 4724 4398 4064	Primer #4 Extension 3558 Da Products ————————————————————————————————————	Mass (Da) 5071 4776 4451 4165
t	3769	 a	3855
nnng nna nc	4724 4398 4064	nnnt nnc na	4776 4451 4165



Mass (daltons)

FIG. 3

Title: USE OF NUCLEOTIDE ANALOSS IN THE ANALYSIS OF OLIGONUCLEOTIDE MIXTURES AND IN HIGHLY MULTIPLEXED NUCLEIC ACID SEQUENCING Docket No.: 25491-2408B Applicant: Cantor acri

Applicant: Cantor et al. Filed: August 20, 2003

FIG. 4a Partially Duplex Hairpin Primer Single-stranded Target IIII TGĂĂŤĠ 3'ACTTACATCTGGAGTGCA5' ANNEAL LIGATE TTGAATG FIG. 4b I I ACTTACATCTGGAGTGCA SEQUENCING REACTION with Mass-matched Terminators (M) FIG. 4c TTGAATGTAGAM ACTTACATCTGGAGTGCA TGAATGTAM ACTTACATCTGGAGTGCA TITGAATGM CTTACATCTGGAGTGCA I I TGAATGTAGM | ACTTACATCTGGAGTGCA ACTTACATCTGGAGTGCA SINGLE STRAND-SPECIFIC **NUCLEASE** FIG. 4d IIIITGÄÄTGTÄGÄM III TGAATGTAGM ACTTACATCT TGAATGTAM TTTGÄÄTGTM ACTTACATO IJACTTACA I I I ŤĠĂĂŤĠ

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Sheet 5 of 8

THIE: USE OF NUCLEOTIDE ANALOGS IN THE ANALYSIS OF
OLIGONUCLEOTIDE MIXTURES AND IN HIGHLY
MULTIPLEXED NUCLEIC ACID SEQUENCING

Docket No.: 25491-24088
Applicant: Cantor et al.
Filed: August 20, 2003

Reaction Products	Mass (Da)
TITTIT GĂĂTGTĂGĂM LIIIIACTTACATCIG	12868.6
TITTIT GAATGTAGM	12227.2
TITIT T G À À T G T À M LILLIA CTTACATO	11594.8
TITTIT GĂĂŤĠŤM LIIIIACTTACAT	10992.4
TITITI T Ğ Á Ä Ť Ġ M LILLIA CTTA CA	10384.0

FIG. 5a

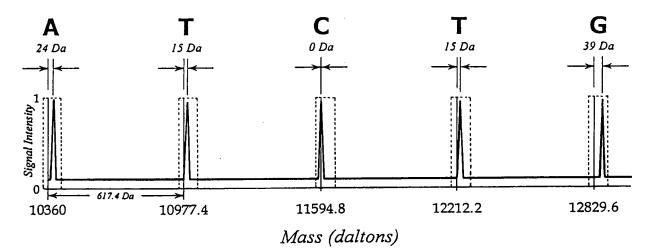
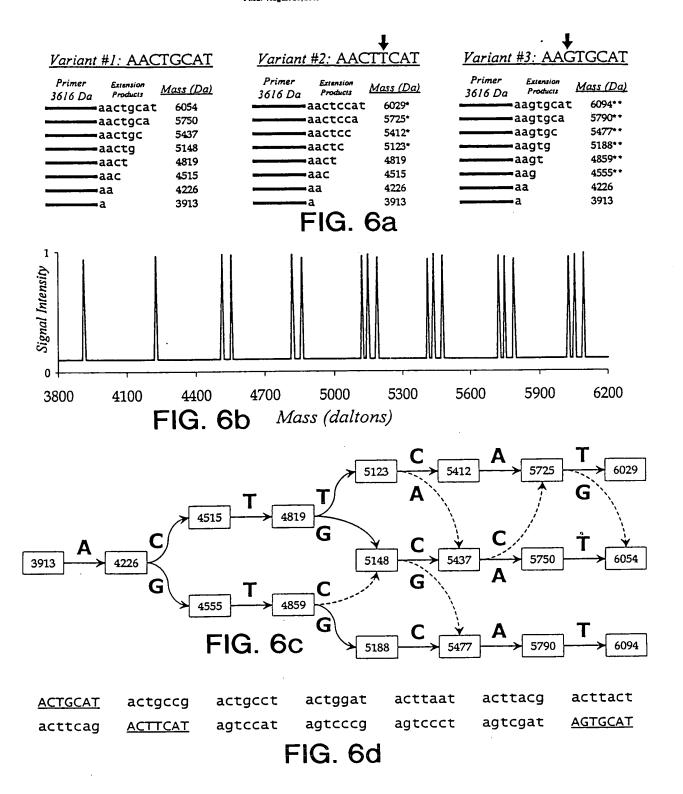


FIG. 5b

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Sheet 6 of 8

Title: USE OF NUCLEOTIDE ANALOGS IN THE ANALYSIS OF
OLICONICLE FOTIDE MISTURES AND IN HIGHLY
MULTIPLEXED NUCLEIC ACID SEQUENCING
Docket No.: 25491-2408B
Applicant: Cantor et al.
Filed: August 20, 2003



HELLER EHRMAN WHITE & MCAULIFFE LLP
Sheet 7 of 8

Title: USE OF NUCLECTIDE ANALOGS IN THE ANALYSIS OF
OLIGONUCLECTIDE MIXTURES AND IN HIGHLY
MULTIPLEXED NUCLEIC ACID SEQUENCING
Docket No.: 25491-2408B

Applicant: Cantor et al. Filed: August 20, 2003

4134

3824

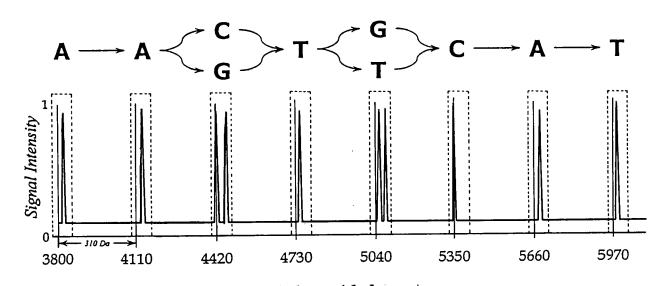
•na

Variant #1: AACI	GCAT	Variant #2: AAC	TTCAT	Variant #3: AA	GTGCAT
Primer Extension A 3527 Da Products A ————————————————————————————————————	5985 5684 5350 5080 4745	Primer Extension 3527 Da Products ————————————————————————————————————	Mass (Da) 5985 5684 5350 5055* 4745 4420	Primer Extension 3527 Da Products	

FIG. 7a

4134

3824



Mass (daltons)

FIG. 7b

$$AA \binom{C}{G} T \binom{G}{T} CAT$$
FIG. 7c

AACTGCAT AACTTCAT AAGTGCAT aagttcat

4134

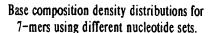
3824

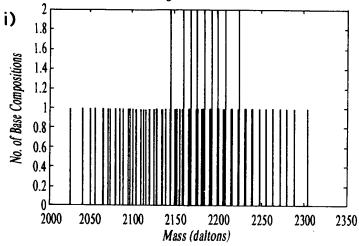
-na

FIG. 7d

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Sheet 8 of 8
USE OF NUCLEOTIDE ANALOGS IN THE ANALYSIS OF
OLIGONUCLEOTIDE MIXTURES AND IN HIGHLY
MULTIPLEXED NUCLEIC ACID SEQUENCING
No.: 25491-2408B

Applicant: Cantor et al. Filed: August 20, 2003





C = 289.2

T = 304.2

A = 313.2

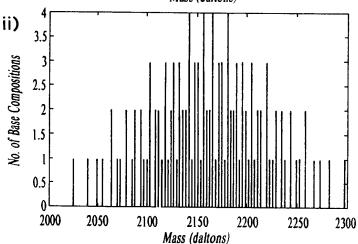
G = 329.2

Naturally Occurring Bases

Peaks can be closer than one dalton

Total No. of different base compositions = 120

Actual number of represented masses = 110 Avg. No. of compositions per mass value = 1.091



C = 289.2

T = 304.2

A = 313.2

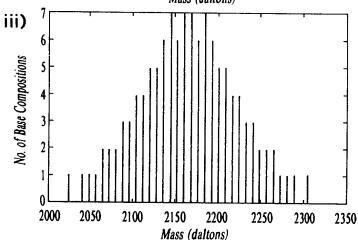
G = 328.2

Substitution with 7-deaza-dG

Minimum peak seperation = 3 daltons Number of allowed mass values = 92

Actual number of represented masses = 64

Avg. No. of compositions per mass value = 1.875



C = 289.2

T = 305.2

A = 313.2

G = 329.2

Substitution with deutero-dT

Minimum peak seperation = 8 daltons Number of allowed mass values = 36

Actual number of represented masses = 34

Avg. No. of compositions per mass value = 3.529

FIG. 8